

SEQUENCE LISTING <110> SYKEN, JOSH MUNGER, KARL <120> METHODS AND REAGENTS TO REGULATE APOPTOSIS <130> HMV-054.01 <140> 09/908,992 <141> 2001-07-19 <150> 60/219,718 <151> 2000-07-19 <150> 60/219,537 <151> 2000-07-20 <160> 27 <170> PatentIn Ver. 2.1 <210> 1 <211> 2656 <212> DNA <213> Homo sapiens <220> <221> CDS <222> (32)..(1471) <400> 1 . gaattegegg eegeagagte eeegggeeaa g atg get geg egg tge tee aca 52 Met Ala Ala Arg Cys Ser Thr cgc tgg ttg ctg gtg gtt gtg ggg acc ccg cgg ctg ccg gct ata tcg Arg Trp Leu Leu Val Val Val Gly Thr Pro Arg Leu Pro Ala Ile Ser 10

100

ggt aga ggg gcc cgg ccc agg gag ggc gtg gtg ggg gca tgg ctg Gly Arg Gly Ala Arg Pro Pro Arg Glu Gly Val Val Gly Ala Trp Leu 25

age ege aag etg age gte eee gee ttt geg tet tee etg ace tet tge Ser Arg Lys Leu Ser Val Pro Ala Phe Ala Ser Ser Leu Thr Ser Cys 40 45

ggc ccc cga gcg ctg ctg aca ttg aga cct ggt gtc agc ctt aca gga 244 Gly Pro Arg Ala Leu Leu Thr Leu Arg Pro Gly Val Ser Leu Thr Gly

aca aaa cat aac cct ttc att tgt act gcc tcc ttc cac acg agt gcc 292 Thr Lys His Asn Pro Phe Ile Cys Thr Ala Ser Phe His Thr Ser Ala 75 80

,

•

			•						2							
			Lys					Gln					Pro		aat Asn	340
		Gln													aag Lys	388
															ttc Phe 135	436
tcc Ser	cag Gln	ctg Leu	gca Ala	gaa Glu 140	gcc Ala	tat Tyr	gag Glu	gtt Val	ttg Leu 145	agt Ser	gat Asp	gag Glu	gtg Val	aag Lys 150	agg Arg	484
				gcc Ala												532
				agc Ser												580
gag Glu	ctg Leu 185	ttc Phe	agg Arg	aag Lys	atc Ile	ttt Phe 190	ggc Gly	gag Glu	ttc Phe	tca Ser	tcc Ser 195	tct Ser	tca Ser	ttt Phe	gga Gly	628
gat Asp 200	ttc Phe	cag Gln	acc Thr	gtg Val	ttt Phe 205	gat Asp	cag Gln	cct Pro	cag Gln	gaa Glu 210	tac Tyr	ttc Phe	atg Met	gag Glu	ttg Leu 215	676
aca Thr	ttc Phe	aat Asn	caa Gln	gct Ala 220	gca Ala	aag Lys	G1 y 999	gtc Val	aac Asn 225	aag Lys	gag Glu	ttc Phe	acc Thr	gtg Val 230	aac Asn	724
atc Ile	atg Met	gac Asp	acg Thr 235	tgt Cys	gag Glu	cgc Arg	tgc Cys	aac Asn 240	ggc Gly	aag Lys	gly aaa	aac Asn	gag Glu 245	ccc Pro	ggc ggc	772
acc Thr	aag Lys	gtg Val 250	cag Gln	cat His	tgc Cys	cac His	tac Tyr 255	tgt Cys	ggc Gly	ggc Gly	tcc Ser	ggc Gly 260	atg Met	gaa Glu	acc Thr	820
atc Ile	aac Asn 265	aca Thr	ggc Gly	cct Pro	ttt Phe	gtg Val 270	atg Met	cgt Arg	tcc Ser	acg Thr	tgt Cys 275	agg Arg	aga Arg	tgt Cys	ggt Gly	868
ggc Gly 280	cgc Arg	ggc Gly	tcc Ser	atc Ile	atc Ile 285	ata Ile	tcg Ser	ccc Pro	tgt Cys	gtg Val 290	gtc Val	tgc Cys	agg Arg	gga Gly	gca Ala 295	916
			Lys	cag Gln 300				Val								964

•																
						•			3							
gto Val	gag Glu	gat Asp	ggc Gly 315	Gln	acc Thr	gtg Val	agg Arg	atg Met 320	cct Pro	gtg Val	gga Gly	aaa Lys	agg Arg 325	gaa Glu	att Ile	1012
tto Phe	att Ile	acg Thr 330	Phe	agg Arg	gtg Val	cag Gln	aaa Lys 335	agc Ser	cct Pro	gtg Val	ttc Phe	cgg Arg 340	agg Arg	gac Asp	ggc Gly	1060
gcā Alā	gac Asp 345	Ile	cac His	tcc Ser	gac Asp	ctc Leu 350	ttt Phe	att Ile	tct Ser	ata Ile	gct Ala 355	cag Gln	gct Ala	ctt Leu	ctt Leu	1108
999 Gly 360	gly,	aca Thr	gcc Ala	aga Arg	gcc Ala 365	cag Gln	ggc Gly	ctg Leu	tac Tyr	gag Glu 370	Thr	atc Ile	aac Asn	gtg Val	acg Thr 375	1156
ato Ile	ccc Pro	cct Pro	Gly aaa	act Thr 380	cag Gln	aca Thr	gac Asp	cag Gln	aag Lys 385	att Ile	cgg Arg	atg Met	ggt Gly	390 Gly 393	aaa Lys	1204
ggc	atc Ile	ccc Pro	cgg Arg 395	att Ile	aac Asn	agc Ser	tac Tyr	ggc Gly 400	tac Tyr	gga Gly	gac Asp	cac His	tac Tyr 405	atc Ile	cac His	1252
atc Ile	aag Lys	ata Ile 410	cga Arg	gtt Val	cca Pro	aag Lys	agg Arg 415	cta Leu	acg Thr	agc Ser	cgg Arg	cag Gln 420	cag Gln	agc Ser	ctg Leu	1300
atc Ile	ctg Leu 425	agc Ser	tac Tyr	gcc Ala	gag Glu	gac Asp 430	gag Glu	aca Thr	gat Asp	gtg Val	gag Glu 435	gly ggg	acg Thr	gtg Val	aac Asn	1348
ggc Gly 440	Val	acc Thr	ctc Leu	acc Thr	agc Ser 445	tct Ser	ggt Gly	ggc Gly	agc Ser	acc Thr 450	atg Met	gat Asp	agc Ser	tcc Ser	gca Ala 455	1396
gga Gly	agc Ser	aag Lys	gct Ala	agg Arg 460	cgt Arg	gag Glu	gct Ala	gly aaa	gag Glu 465	gac Asp	gag Glu	gag Glu	gga Gly	ttc Phe 470	ctt Leu	1444
tcc Ser	aaa Lys	ctt Leu	aag Lys 475	aaa Lys	atg Met	ttt Phe	acc Thr	tca Ser 480	tgat	atco	ca g	rccga	ggaa	a		1491
aag	atcca	act g	gaaa	.ctag	g cc	ggga	.agca	ı gca	gccc	ctc	caag	iggcc	ag ,g	gcac	ctggg	1551
agad	ggga	igg a	ttcc	agaa	c ag	cago	actg	g agc	tccc	acc	cgca	gagc	ct c	tgga	.cggcc	1611
ttg	gcaac	ag c	aaaa	tcat	a aa	acaa	cacc	tct	ctcc	acg	gaaa	ggtc	ac a	gtgg	acagc	1671
ccg	ggcag	jta g	gatg	cagc	c cc	agag	gctg	gtg	gcag	ttt	cctg	tcca	tt g	gtag	gtgac	1731
															gaatt	
															taagc	
tgca	itcaa	gt t	acga	agtg	a tt	aatt	tcct	tct	cagc	aaa	cctc	cggg	ag g	ttcc	agaat	1911

gagttettee tgacaggttg tetteactgg gagegtgggg eccecaggee ecaceageae 1971
egtecteece taatgagggg ecctgeegag geateagetg etetgeteag ttagttttta 2031
tteeeggggt accaageage tgeacagteg gtgeetggga ageaegttaa aggeeeagag 2091
agateettggg ggttetgete tgacegtgg ggtggtgate ettgteagga tgtacagtee 2151
ttgeteecae eccateeggg atggeegeet gteeetgaet attgagteet gttgttgtaa 2211
geeaggeatg gagggeteet geeettetge tgageeacag eccattgeag eactgttgge 2271
geeagaette agetgeettg ggaactgaag ecctgeeaet gttgetagte aggggettgg 2331
tteeteecaet tacactgttg acatetatt tetgaagtgt gtttaaatta tteagtgeta 2391
atcattgttt ttteetttgt aaatgstgat teagaaaagg aaageacagg etaageagtt 2451
gaaggtteee eaceatteag tgagageaga acceeeatte eccageetet getggtagea 2571
eactteteaa eagttteett tetgtttee tttataatte actaaaataa ageatetatt 2631
agtgtetgaa aaaaaaaaa aaaaa 2656

<210> 2 <211> 1443 <212> DNA <213> Homo sapiens <220> <221> CDS <222> (1)..(1440) <400> 2 atg gct gcg cgg tgc tcc aca cgc tgg ttg ctg gtg gtt gtg ggg acc 48 Met Ala Ala Arg Cys Ser Thr Arg Trp Leu Leu Val Val Gly Thr ccg cgg ctg ccg gct ata tcg ggt aga ggg gcc cgg ccg ccc agg gag 96 Pro Arg Leu Pro Ala Ile Ser Gly Arg Gly Ala Arg Pro Pro Arg Glu ggc gtg gtg ggg gca tgg ctg agc cgc aag ctg agc gtc ccc gcc ttt 144 Gly Val Val Gly Ala Trp Leu Ser Arg Lys Leu Ser Val Pro Ala Phe geg tet tee etg ace tet tge gge eee ega geg etg etg aca ttg aga 192 Ala Ser Ser Leu Thr Ser Cys Gly Pro Arg Ala Leu Leu Thr Leu Arg cet ggt gte age ett aca gga aca aaa cat aac eet tte att tgt act 240 Pro Gly Val Ser Leu Thr Gly Thr Lys His Asn Pro Phe Ile Cys Thr 70 75 80

							5						
_			_	_	_	_	_		_	_		cag Gln	288
					aat Asn								336
					aag Lys								384
					ttc Phe 135								432
					agg Arg								480
					agc Ser								528
					gag Glu								576
					gga Gly								624
					ttg Leu 215								672
					aac Asn								720
	_				ggc Gly	_		_		-		_	768
					acc Thr								816
					ggt Gly								864
					gca Ala 295								912

atg Met 305	Ile	cct Pro	gtg Val	cct Pro	gca Ala 310	gga Gly	gtc Val	gag Glu	gat Asp	ggc Gly 315	cag Gln	acc Thr	gtg Val	agg Arg	atg Met 320	960
cct Pro	gtg Val	gga Gly	aaa Lys	agg Arg 325	gaa Glu	att Ile	ttc Phe	att Ile	acg Thr 330	ttc Phe	agg Arg	gtg Val	cag Gln	aaa Lys 335	agc Ser	1008
					gac Asp											1056
tct Ser	ata Ile	gct Ala 355	cag Gln	gct Ala	ctt Leu	ctt Leu	360 999	gga Gly	aca Thr	gcc Ala	aga Arg	gcc Ala 365	cag Gln	ggc	ctg Leu	1104
tac Tyr	gag Glu 370	acg Thr	atc Ile	aac Asn	gtg Val	acg Thr 375	atc Ile	ccc Pro	cct Pro	elà aaa	act Thr 380	cag Gln	aca Thr	gac Asp	cag Gln	1152
aag Lys 385	att Ile	cgg Arg	atg Met	ggt Gly	390 399	aaa Lys	ggc Gly	atc Ile	ccc Pro	cgg Arg 395	att Ile	aac Asn	agc Ser	tac Tyr	ggc Gly 400	1200
tac Tyr	gga Gly	gac Asp	cac His	tac Tyr 405	atc Ile	cac His	atc Ile	aag Lys	ata Ile 410	cga Arg	gtt Val	cca Pro	aag Lys	agg Arg 415	cta Leu	1248
acg Thr	agc Ser	cgg Arg	cag Gln 420	cag Gln	agc Ser	ctg Leu	atc Ile	ctg Leu 425	agc Ser	tac Tyr	gcc Ala	gag Glu	gac Asp 430	gag Glu	aca Thr	1296
gat Asp	gtg Val	gag Glu 435	gly aaa	acg Thr	gtg Val	aac Asn	ggc Gly 440	gtc Val	acc Thr	ctc Leu	acc Thr	agc Ser 445	tct Ser	ggt Gly	ggc Gly	1344
agc Ser	acc Thr 450	atg Met	gat Asp	agc Ser	tcc Ser	gca Ala 455	gga Gly	agc Ser	aag Lys	gct Ala	agg Arg 460	cgt Arg	gag Glu	gct Ala	999 999	1392
gag Glu 465	gac Asp	gag Glu	gag Glu	gga Gly	ttc Phe 470	ctt Leu	tcc Ser	aaa Lys	ctt Leu	aag Lys 475	aaa Lys	atg Met	ttt Phe	acc Thr	tca Ser 480	1440
tga																1443

<210> 3 <211> 1362

<212> DNA

<213> Homo sapiens

<220>

<221> CDS

<222> (1)..(1359)

<40	0 > 3																	
	Ala				Ser		cgc Arg			Leu						48		
ccg Pro	cgg Arg	ctg Leu	ccg Pro 20	Ala	ata Ile	tcg Ser	ggt Gly	aga Arg 25	gjà aaa	gcc Ala	cgg Arg	ccg Pro	ccc Pro 30	agg Arg	gag Glu	96		
			Gly				agc Ser 40									144		
							ggc Gly									192		
cct Pro 65	Gly	gtc Val	agc Ser	ctt Leu	aca Thr 70	gga Gly	aca Thr	aaa Lys	cat His	aac Asn 75	cct Pro	ttc Phe	att Ile	tgt Cys	act Thr 80	240		
gcc Ala	tcc Ser	ttc Phe	cac His	acg Thr 85	agt Ser	gcc Ala	cct Pro	ttg Leu	gcc Ala 90	aaa Lys	gaa Glu	gat Asp	tat Tyr	tat Tyr 95	cag Gln	288		
							gcc Ala									336		
tat Tyr	tat Tyr	cag Gln 115	ctt Leu	gcc Ala	aag Lys	aag Lys	tat Tyr 120	cac His	cct Pro	gac Asp	aca Thr	aat Asn 125	aag Lys	gat Asp	gat Asp	384	٠	
ccc Pro	aaa Lys 130	gcc Ala	aag Lys	gag Glu	aag Lys	ttc Phe 135	tcc Ser	cag Gln	ctg Leu	gca Ala	gaa Glu 140	gcc Ala	tat Tyr	gag Glu	gtt Val	432		
ttg Leu 145	agt Ser	gat Asp	gag Glu	gtg Val	aag Lys 150	agg Arg	aag Lys	cag Gln	tac Tyr	gat Asp 155	gcc Ala	tac Tyr	ggc Gly	tct Ser	gca Ala 160	480		
ggc Gly	ttc Phe	gat Asp	cct Pro	165 Gly 999	gcc Ala	agc Ser	ggc Gly	tcc Ser	cag Gln 170	cat His	agc Ser	tac Tyr	tgg Trp	aag Lys 175	gga Gly	528		
ggc Gly	ccc Pro	act Thr	gtg Val 180	gac Asp	ccc Pro	gag Glu	gag Glu	ctg Leu 185	ttc Phe	agg Arg	aag Lys	atc Ile	ttt Phe 190	ggc	gag Glu	576		
ttc Phe	Ser	tcc Ser 195	tct Ser	tca Ser	ttt Phe	gga Gly	gat Asp 200	ttc Phe	cag Gln	acc Thr	gtg Val	ttt Phe 205	gat Asp	cag Gln	cct Pro	624		
cag Gln	gaa Glu 210	tac Tyr	ttc Phe	atg Met	Glu	ttg Leu 215	aca Thr	ttc Phe	aat Asn	Gln	gct Ala 220	gca Ala	aag Lys	Gly aaa	gtc Val	672		

		ttc Phe										720
		aac Asn										768
		ggc Gly 260										816
		agg Arg										864
		tgc Cys										912
		gtg Val										960
		aaa Lys										1008
		cgg Arg 340										1056
	_	cag Gln	_		 	_	_	_	_		_	1104
	_	atc Ile		 _				_		_	_	1152
		atg Met										1200
		cac His										1248
		cag Gln 420										1296
		Gly ggg										1344

```
aga tcc act gga aac tag
                                                                  1362
Arg Ser Thr Gly Asn
    450
<210> 4
<211> 1242
<212> DNA
<213> Homo sapiens
<400> 4
gtcagcctta caggaacaaa acataaccct ttcatttgta ctgcctcctt ccacacgagt 60
gcccctttgg ccaaagaaga ttattatcag atattaggag tgcctcgaaa tgccagccag 120
aaagagatca agaaagccta ttatcagctt gccaagaagt atcaccctga cacaaataag 180
gatgatccca aagccaagga gaagttctcc cagctggcag aagcctatga ggttttgagt 240
gatgaggtga agaggaagca gtacgatgcc tacggctctg caggcttcga tcctggggcc 300
ageggetece ageatageta etggaaggga ggeeceaetg tggaeceega ggagetgtte 360
aggaagatet tiggegagit eteateetet teatitiggag atticeagae egigtitigat 420
cagcctcagg aatacttcat ggagttgaca ttcaatcaag ctgcaaaggg ggtcaacaag 480
gagttcaccg tgaacatcat ggacacgtgt gagcgctgca acggcaaggg gaacgagccc 540
ggcaccaagg tgcagcattg ccactactgt ggcggctccg gcatggaaac catcaacaca 600
ggcccttttg tgatgcgttc cacgtgtagg agatgtggtg gccgcggctc catcatcata 660
tegecetgtg tggtetgeag gggageagga caageeaage agaaaaageg agtgatgate 720
cctgtgcctg caggagtcga ggatggccag accgtgagga tgcctgtggg aaaaaagggaa 780
attttcatta cgttcagggt gcagaaaagc cctgtgttcc ggagggacgg cgcagacatc 840
cactccgacc tetttattte tatageteag getettettg ggggaacage cagageccag 900
ggcctgtacg agacgatcaa cgtgacgatc ccccctggga ctcagacaga ccagaagatt 960
cggatgggtg ggaaaggcat cccccggatt aacagctacg gctacggaga ccactacatc 1020
cacatcaaga tacgagttcc aaagaggcta acgagccggc agcagagcct gatcctgagc 1080
tacgccgagg acgagacaga tgtggagggg acggtgaacg gcgtcaccct caccagctct 1140
ggtggcagca ccatggatag ctccgcagga agcaaggcta ggcgtgaggc tggggaggac 1200
gaggagggat tcctttccaa acttaagaaa atgtttacct ca
<210> 5
<211> 1161
<212> DNA
<213> Homo sapiens
<400> 5
gtcagcctta caggaacaaa acataaccct ttcatttgta ctgcctcctt ccacacgagt 60
gcccctttgg ccaaagaaga ttattatcag atattaggag tgcctcgaaa tgccagccag 120
aaagagatca agaaagccta ttatcagctt gccaagaagt atcaccctga cacaaataag 180
gatgatccca aagccaagga gaagttctcc cagctggcag aagcctatga qqttttqaqt 240
gatgaggtga agaggaagca gtacgatgcc tacggctctg caqqcttcqa tcctqqqqcc 300
ageggeteee ageatageta etggaaggga ggeeceaetg tggaeceega ggagetgtte 360
aggaagatet ttggegagtt eteateetet teatttggag attteeagae egtgtttgat 420
cagcctcagg aatacttcat ggagttgaca ttcaatcaag ctgcaaaggg ggtcaacaag 480
gagttcaccg tgaacatcat ggacacgtgt gagcgctgca acggcaaggg gaacgagccc 540
ggcaccaagg tgcagcattg ccactactgt ggcggctccg gcatggaaac catcaacaca 600
ggcccttttg tgatgcgttc cacgtgtagg agatgtggtg gccgcggctc catcatcata 660
cctgtgcctg caggagtcga ggatggccag accgtgagga tgcctgtggg aaaaagggaa 780
attttcatta cgttcagggt gcagaaaagc cctgtgttcc ggagggacgg cgcagacatc 840
cactccgacc tetttattte tatageteag getettettg ggggaacage cagageecag 900
ggcctgtacg agacgatcaa cgtgacgatc ccccctggga ctcagacaga ccagaagatt 960
cggatgggtg ggaaaggcat cccccggatt aacagctacg gctacggaga ccactacatc 1020
cacatcaaga tacgagttcc aaagaggcta acgagccggc agcagagcct gatcctgagc 1080
```

tacgccgagg acgagacaga tgtggagggg acggtgaacg gcgtcaccct caccagctct 1140 ggaaaaagat ccactqqaaa c <210> 6 <211> 99 <212> DNA <213> Homo sapiens <400> 6 ggcagcacca tggatagctc cgcaggaagc aaggctaggc gtgaggctgg ggaggacgag 60 gagggattcc tttccaaact taagaaaatg tttacctca <210> 7 <211> 18 <212> DNA <213> Homo sapiens <400> 7 aaaagatcca ctggaaac 18 <210> 8 <211> 480 <212> PRT <213> Homo sapiens <400> 8 Met Ala Ala Arg Cys Ser Thr Arg Trp Leu Leu Val Val Gly Thr 10 Pro Arg Leu Pro Ala Ile Ser Gly Arg Gly Ala Arg Pro Pro Arg Glu Gly Val Val Gly Ala Trp Leu Ser Arg Lys Leu Ser Val Pro Ala Phe 40 Ala Ser Ser Leu Thr Ser Cys Gly Pro Arg Ala Leu Leu Thr Leu Arg Pro Gly Val Ser Leu Thr Gly Thr Lys His Asn Pro Phe Ile Cys Thr 70 Ala Ser Phe His Thr Ser Ala Pro Leu Ala Lys Glu Asp Tyr Tyr Gln 90 Ile Leu Gly Val Pro Arg Asn Ala Ser Gln Lys Glu Ile Lys Lys Ala 100 Tyr Tyr Gln Leu Ala Lys Lys Tyr His Pro Asp Thr Asn Lys Asp Asp 120 Pro Lys Ala Lys Glu Lys Phe Ser Gln Leu Ala Glu Ala Tyr Glu Val 130 135 Leu Ser Asp Glu Val Lys Arg Lys Gln Tyr Asp Ala Tyr Gly Ser Ala 150 155

160

- Gly Phe Asp Pro Gly Ala Ser Gly Ser Gln His Ser Tyr Trp Lys Gly
 165 170 175
- Gly Pro Thr Val Asp Pro Glu Glu Leu Phe Arg Lys Ile Phe Gly Glu 180 185 190
- Phe Ser Ser Ser Phe Gly Asp Phe Gln Thr Val Phe Asp Gln Pro 195 200 205
- Gln Glu Tyr Phe Met Glu Leu Thr Phe Asn Gln Ala Ala Lys Gly Val 210 215 220
- Asn Lys Glu Phe Thr Val Asn Ile Met Asp Thr Cys Glu Arg Cys Asn 225 235 240
- Gly Lys Gly Asn Glu Pro Gly Thr Lys Val Gln His Cys His Tyr Cys 245 250 255
- Gly Gly Ser Gly Met Glu Thr Ile Asn Thr Gly Pro Phe Val Met Arg 260 265 270
- Ser Thr Cys Arg Arg Cys Gly Gly Arg Gly Ser Ile Ile Ile Ser Pro 275 280 285
- Cys Val Val Cys Arg Gly Ala Gly Gln Ala Lys Gln Lys Lys Arg Val 290 295 300
- Met Ile Pro Val Pro Ala Gly Val Glu Asp Gly Gln Thr Val Arg Met 305 310 315 320
- Pro Val Gly Lys Arg Glu Ile Phe Ile Thr Phe Arg Val Gln Lys Ser 325 330 335
- Pro Val Phe Arg Arg Asp Gly Ala Asp Ile His Ser Asp Leu Phe Ile 340 345 350
- Ser Ile Ala Gln Ala Leu Leu Gly Gly Thr Ala Arg Ala Gln Gly Leu 355 360 365
- Tyr Glu Thr Ile Asn Val Thr Ile Pro Pro Gly Thr Gln Thr Asp Gln 370 375 380
- Lys Ile Arg Met Gly Gly Lys Gly Ile Pro Arg Ile Asn Ser Tyr Gly 385 390 395 400
- Tyr Gly Asp His Tyr Ile His Ile Lys Ile Arg Val Pro Lys Arg Leu 405 410 415
- Thr Ser Arg Gln Gln Ser Leu Ile Leu Ser Tyr Ala Glu Asp Glu Thr 420 425 430
- Asp Val Glu Gly Thr Val Asn Gly Val Thr Leu Thr Ser Ser Gly Gly
 435 440 445
- Ser Thr Met Asp Ser Ser Ala Gly Ser Lys Ala Arg Arg Glu Ala Gly 450 455 460

Glu Asp Glu Glu Gly Phe Leu Ser Lys Leu Lys Lys Met Phe Thr Ser 465 470 475 480

<210> 9

<211> 453

<212> PRT

<213> Homo sapiens

<400> 9

Met Ala Ala Arg Cys Ser Thr Arg Trp Leu Leu Val Val Val Gly Thr
1 5 10 15

Pro Arg Leu Pro Ala Ile Ser Gly Arg Gly Ala Arg Pro Pro Arg Glu 20 25 30

Gly Val Val Gly Ala Trp Leu Ser Arg Lys Leu Ser Val Pro Ala Phe 35 40 45

Ala Ser Ser Leu Thr Ser Cys Gly Pro Arg Ala Leu Leu Thr Leu Arg
50 55 60

Pro Gly Val Ser Leu Thr Gly Thr Lys His Asn Pro Phe Ile Cys Thr 65 70 75 80

Ala Ser Phe His Thr Ser Ala Pro Leu Ala Lys Glu Asp Tyr Tyr Gln
85 90 95

Ile Leu Gly Val Pro Arg Asn Ala Ser Gln Lys Glu Ile Lys Lys Ala 100 105 110

Tyr Tyr Gln Leu Ala Lys Lys Tyr His Pro Asp Thr Asn Lys Asp Asp 115 120 125

Pro Lys Ala Lys Glu Lys Phe Ser Gln Leu Ala Glu Ala Tyr Glu Val 130 135 140

Leu Ser Asp Glu Val Lys Arg Lys Gln Tyr Asp Ala Tyr Gly Ser Ala 145 150 155 160

Gly Phe Asp Pro Gly Ala Ser Gly Ser Gln His Ser Tyr Trp Lys Gly
165 170 175

Gly Pro Thr Val Asp Pro Glu Glu Leu Phe Arg Lys Ile Phe Gly Glu 180 185 190

Phe Ser Ser Ser Phe Gly Asp Phe Gln Thr Val Phe Asp Gln Pro 195 200 205

Gln Glu Tyr Phe Met Glu Leu Thr Phe Asn Gln Ala Ala Lys Gly Val 210 215 220

Asn Lys Glu Phe Thr Val Asn Ile Met Asp Thr Cys Glu Arg Cys Asn 225 230 235 240

Gly Lys Gly Asn Glu Pro Gly Thr Lys Val Gln His Cys His Tyr Cys 245 250 255 Gly Gly Ser Gly Met Glu Thr Ile Asn Thr Gly Pro Phe Val Met Arg
260 265 270

Ser Thr Cys Arg Arg Cys Gly Gly Arg Gly Ser Ile Ile Ile Ser Pro 275 280 285

Cys Val Val Cys Arg Gly Ala Gly Gln Ala Lys Gln Lys Lys Arg Val 290 295 300

Met Ile Pro Val Pro Ala Gly Val Glu Asp Gly Gln Thr Val Arg Met 305 310 315 320

Pro Val Gly Lys Arg Glu Ile Phe Ile Thr Phe Arg Val Gln Lys Ser 325 330 335

Pro Val Phe Arg Arg Asp Gly Ala Asp Ile His Ser Asp Leu Phe Ile 340 345 350

Ser Ile Ala Gln Ala Leu Leu Gly Gly Thr Ala Arg Ala Gln Gly Leu 355 360 365

Tyr Glu Thr Ile Asn Val Thr Ile Pro Pro Gly Thr Gln Thr Asp Gln 370 375 380

Lys Ile Arg Met Gly Gly Lys Gly Ile Pro Arg Ile Asn Ser Tyr Gly 385 390 395 400

Tyr Gly Asp His Tyr Ile His Ile Lys Ile Arg Val Pro Lys Arg Leu 405 410 415

Thr Ser Arg Gln Gln Ser Leu Ile Leu Ser Tyr Ala Glu Asp Glu Thr 420 425 430

Asp Val Glu Gly Thr Val Asn Gly Val Thr Leu Thr Ser Ser Gly Lys
435
440
445

Arg Ser Thr Gly Asn 450

<210> 10

<211> 414

<212> PRT

<213> Homo sapiens

<400> 10

Val Ser Leu Thr Gly Thr Lys His Asn Pro Phe Ile Cys Thr Ala Ser 1 5 10 15

Phe His Thr Ser Ala Pro Leu Ala Lys Glu Asp Tyr Tyr Gln Ile Leu
20 25 30

Gly Val Pro Arg Asn Ala Ser Gln Lys Glu Ile Lys Lys Ala Tyr Tyr

Gln Leu Ala Lys Lys Tyr His Pro Asp Thr Asn Lys Asp Asp Pro Lys
50 60

- Ala Lys Glu Lys Phe Ser Gln Leu Ala Glu Ala Tyr Glu Val Leu Ser 65 70 75 80
- Asp Glu Val Lys Arg Lys Gln Tyr Asp Ala Tyr Gly Ser Ala Gly Phe 85 90 95
- Asp Pro Gly Ala Ser Gly Ser Gln His Ser Tyr Trp Lys Gly Gly Pro
 100 105 110
- Thr Val Asp Pro Glu Glu Leu Phe Arg Lys Ile Phe Gly Glu Phe Ser 115 120 125
- Ser Ser Ser Phe Gly Asp Phe Gln Thr Val Phe Asp Gln Pro Gln Glu
 130 135 140
- Tyr Phe Met Glu Leu Thr Phe Asn Gln Ala Ala Lys Gly Val Asn Lys 145 150 155 160
- Glu Phe Thr Val Asn Ile Met Asp Thr Cys Glu Arg Cys Asn Gly Lys 165 170 175
- Gly Asn Glu Pro Gly Thr Lys Val Gln His Cys His Tyr Cys Gly Gly
 180 185 190
- Ser Gly Met Glu Thr Ile Asn Thr Gly Pro Phe Val Met Arg Ser Thr 195 200 205
- Cys Arg Arg Cys Gly Gly Arg Gly Ser Ile Ile Ile Ser Pro Cys Val 210 215 220
- Val Cys Arg Gly Ala Gly Gln Ala Lys Gln Lys Lys Arg Val Met Ile 225 230 235 240
- Pro Val Pro Ala Gly Val Glu Asp Gly Gln Thr Val Arg Met Pro Val 245 250 255
- Gly Lys Arg Glu Ile Phe Ile Thr Phe Arg Val Gln Lys Ser Pro Val 260 265 270
- Phe Arg Arg Asp Gly Ala Asp Ile His Ser Asp Leu Phe Ile Ser Ile 275 280 285
- Ala Gln Ala Leu Leu Gly Gly Thr Ala Arg Ala Gln Gly Leu Tyr Glu 290 295 300
- Thr Ile Asn Val Thr Ile Pro Pro Gly Thr Gln Thr Asp Gln Lys Ile 305 310 315 320
- Arg Met Gly Gly Lys Gly Ile Pro Arg Ile Asn Ser Tyr Gly Tyr Gly 325 330 335
- Asp His Tyr Ile His Ile Lys Ile Arg Val Pro Lys Arg Leu Thr Ser 340 345 350
- Arg Gln Gln Ser Leu Ile Leu Ser Tyr Ala Glu Asp Glu Thr Asp Val 355 360 365

Glu Gly Thr Val Asn Gly Val Thr Leu Thr Ser Ser Gly Gly Ser Thr 370 375 380

Met Asp Ser Ser Ala Gly Ser Lys Ala Arg Arg Glu Ala Gly Glu Asp 385 390 395 400

Glu Glu Gly Phe Leu Ser Lys Leu Lys Lys Met Phe Thr Ser 405 410

<210> 11

<211> 387

<212> PRT

<213> Homo sapiens

<400> 11

Val Ser Leu Thr Gly Thr Lys His Asn Pro Phe Ile Cys Thr Ala Ser 1 5 10 15

Phe His Thr Ser Ala Pro Leu Ala Lys Glu Asp Tyr Tyr Gln Ile Leu 20 25 30

Gly Val Pro Arg Asn Ala Ser Gln Lys Glu Ile Lys Lys Ala Tyr Tyr 35 40 45

Gln Leu Ala Lys Lys Tyr His Pro Asp Thr Asn Lys Asp Asp Pro Lys
50 55 60

Ala Lys Glu Lys Phe Ser Gln Leu Ala Glu Ala Tyr Glu Val Leu Ser 65 70 75 80

Asp Glu Val Lys Arg Lys Gln Tyr Asp Ala Tyr Gly Ser Ala Gly Phe
85 90 95

Asp Pro Gly Ala Ser Gly Ser Gln His Ser Tyr Trp Lys Gly Gly Pro 100 105 110

Thr Val Asp Pro Glu Glu Leu Phe Arg Lys Ile Phe Gly Glu Phe Ser 115 120 125

Ser Ser Ser Phe Gly Asp Phe Gln Thr Val Phe Asp Gln Pro Gln Glu 130 135 140

Tyr Phe Met Glu Leu Thr Phe Asn Gln Ala Ala Lys Gly Val Asn Lys 145 150 155 160

Glu Phe Thr Val Asn Ile Met Asp Thr Cys Glu Arg Cys Asn Gly Lys
165 170 175

Gly Asn Glu Pro Gly Thr Lys Val Gln His Cys His Tyr Cys Gly Gly 180 185 190

Ser Gly Met Glu Thr Ile Asn Thr Gly Pro Phe Val Met Arg Ser Thr 195 200 205

Cys Arg Arg Cys Gly Gly Arg Gly Ser Ile Ile Ile Ser Pro Cys Val 210 215 220 Val Cys Arg Gly Ala Gly Gln Ala Lys Gln Lys Lys Arg Val Met Ile 225 230 235 240

Pro Val Pro Ala Gly Val Glu Asp Gly Gln Thr Val Arg Met Pro Val 245 250 255

Gly Lys Arg Glu Ile Phe Ile Thr Phe Arg Val Gln Lys Ser Pro Val 260 265 270

Phe Arg Arg Asp Gly Ala Asp Ile His Ser Asp Leu Phe Ile Ser Ile 275 280 285

Ala Gln Ala Leu Leu Gly Gly Thr Ala Arg Ala Gln Gly Leu Tyr Glu 290 295 300

Thr Ile Asn Val Thr Ile Pro Pro Gly Thr Gln Thr Asp Gln Lys Ile 305 310 315 320

Arg Met Gly Gly Lys Gly Ile Pro Arg Ile Asn Ser Tyr Gly Tyr Gly 325 330 335

Asp His Tyr Ile His Ile Lys Ile Arg Val Pro Lys Arg Leu Thr Ser 340 345 350

Arg Gln Gln Ser Leu Ile Leu Ser Tyr Ala Glu Asp Glu Thr Asp Val 355 360 365

Glu Gly Thr Val Asn Gly Val Thr Leu Thr Ser Ser Gly Lys Arg Ser 370 380

Thr Gly Asn 385

<210> 12

<211> 480

<212> PRT

<213> Homo sapiens

<400> 12

Met Ala Ala Arg Cys Ser Thr Arg Trp Leu Leu Val Val Gly Thr 1 5 10 15

Pro Arg Leu Pro Ala Ile Ser Gly Arg Gly Ala Arg Pro Pro Arg Glu 20 25 30

Gly Val Val Gly Ala Trp Leu Ser Arg Lys Leu Ser Val Pro Ala Phe 35 . 40

Ala Ser Ser Leu Thr Ser Cys Gly Pro Arg Ala Leu Leu Thr Leu Arg
50 55 60

Pro Gly Val Ser Leu Thr Gly Thr Lys His Asn Pro Phe Ile Cys Thr 65 70 75 80

Ala Ser Phe His Thr Ser Ala Pro Leu Ala Lys Glu Asp Tyr Tyr Gln 85 90 95

- Ile Leu Gly Val Pro Arg Asn Ala Ser Gln Lys Glu Ile Lys Lys Ala 100 105 110
- Tyr Tyr Gln Leu Ala Lys Lys Tyr His Pro Asp Thr Asn Lys Asp Asp 115 120 125
- Pro Lys Ala Lys Glu Lys Phe Ser Gln Leu Ala Glu Ala Tyr Glu Val 130 135 140
- Leu Ser Asp Glu Val Lys Arg Lys Gln Tyr Asp Ala Tyr Gly Ser Ala 145 150 155 160
- Gly Phe Asp Pro Gly Ala Ser Gly Ser Gln His Ser Tyr Trp Lys Gly
 165 170 175
- Gly Pro Thr Val Asp Pro Glu Glu Leu Phe Arg Lys Ile Phe Gly Glu 180 185 190
- Phe Ser Ser Ser Phe Gly Asp Phe Gln Thr Val Phe Asp Gln Pro 195 200 205
- Gln Glu Tyr Phe Met Glu Leu Thr Phe Asn Gln Ala Ala Lys Gly Val 210 215 220
- Asn Lys Glu Phe Thr Val Asn Ile Met Asp Thr Cys Glu Arg Cys Asn 225 230 235 240
- Gly Lys Gly Asn Glu Pro Gly Thr Lys Val Gln His Cys His Tyr Cys 245 250 255
- Gly Gly Ser Gly Met Glu Thr Ile Asn Thr Gly Pro Phe Val Met Arg
 260 265 270
- Ser Thr Cys Arg Arg Cys Gly Gly Arg Gly Ser Ile Ile Ile Ser Pro 275 280 285
- Cys Val Val Cys Arg Gly Ala Gly Gln Ala Lys Gln Lys Lys Arg Val 290 295 300
- Met Ile Pro Val Pro Ala Gly Val Glu Asp Gly Gln Thr Val Arg Met 305 310 315 320
- Pro Val Gly Lys Arg Glu Ile Phe Ile Thr Phe Arg Val Gln Lys Ser 325 330 335
- Pro Val Phe Arg Arg Asp Gly Ala Asp Ile His Ser Asp Leu Phe Ile 340 345 350
- Ser Ile Ala Gln Ala Leu Leu Gly Gly Thr Ala Arg Ala Gln Gly Leu 355 360 365
- Tyr Glu Thr Ile Asn Val Thr Ile Pro Pro Gly Thr Gln Thr Asp Gln 370 375 380
- Lys Ile Arg Met Gly Gly Lys Gly Ile Pro Arg Ile Asn Ser Tyr Gly 385 390 395 400

Tyr Gly Asp His Tyr Ile His Ile Lys Ile Arg Val Pro Lys Arg Leu 405 410 415

Thr Ser Arg Gln Gln Ser Leu Ile Leu Ser Tyr Ala Glu Asp Glu Thr 420 425 430

Asp Val Glu Gly Thr Val Asn Gly Val Thr Leu Thr Ser Ser Gly Gly
435 440 445

Ser Thr Met Asp Ser Ser Ala Gly Ser Lys Ala Arg Arg Glu Ala Gly 450 460

Glu Asp Glu Glu Gly Phe Leu Ser Lys Leu Lys Lys Met Phe Thr Ser 465 470 475 480

<210> 13

<211> 33

<212> PRT

<213> Homo sapiens

<400> 13

Gly Ser Thr Met Asp Ser Ser Ala Gly Ser Lys Ala Arg Arg Glu Ala 1 5 10 15

Gly Glu Asp Glu Gly Phe Leu Ser Lys Leu Lys Lys Met Phe Thr 20 25 30

Ser

<210> 14

<211> 6

<212> PRT

<213> Homo sapiens

<400> 14

Lys Arg Ser Thr Gly Asn

<210> 15

<211> 26

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic peptide

<400> 15

Cys Phe Ile Thr Lys Ala Leu Gly Ile Ser Tyr Gly Arg Lys Lys Arg
1 5 10 15

Arg Gln Arg Arg Pro Pro Gln Gly Ser

```
<210> 16
<211> 12
<212> PRT
<213> Unknown Organism
<220>
<223> Description of Unknown Organism: EGF derived
     peptide
<400> 16
Cys Met His Ile Glu Ser Leu Asp Ser Tyr Thr Cys
<210> 17
<211> 12
<212> PRT
<213> Unknown Organism
<220>
<223> Description of Unknown Organism: EGF derived
      peptide
<400> 17
Cys Met Tyr Ile Glu Ala Leu Asp Lys Tyr Ala Cys
<210> 18
<211> 29
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic
      internalizing peptide
Glu Ala Ala Leu Ala Glu Ala Leu Ala Glu Ala Leu Ala Glu Ala Leu
Ala Glu Ala Leu Ala Glu Ala Leu Glu Ala Leu Ala Ala
             20
                                 25
<210> 19
<211> 8
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Illustrative
      peptide
<400> 19
Gly Asn Ala Ala Ala Arg Arg
                  5
  1
```

20

```
<210> 20
<211> 20
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: Primer
<400> 20
                                                                    20
cgagacagat gtggagggga
<210> 21
<211> 18
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: Primer
<400> 21
                                                                    18
gaataattta aacacact
<210> 22
<211> 36
<212> PRT
<213> Homo sapiens
<400> 22
Ser Ser Gly Gly Ser Thr Met Asp Ser Ser Ala Gly Ser Lys Ala Arg
Arg Glu Ala Gly Glu Asp Glu Glu Gly Phe Leu Ser Lys Leu Lys Lys
                                  25
Met Phe Thr Ser
        35
<210> 23
<211> 9
<212> PRT
<213> Homo sapiens
Ser Ser Gly Lys Arg Ser Thr Gly Asn
                  5
<210> 24
<211> 33
<212> PRT
<213> Homo sapiens
```

```
21
<400> 24
Gly Ser Thr Met Asp Ser Ser Ala Gly Ser Lys Ala Arg Arg Glu Ala
Gly Glu Asp Glu Glu Gly Phe Leu Ser Lys Leu Lys Lys Met Phe Thr
Ser
```

<210> 25 <211> 33 <212> PRT <213> Mus sp. <400> 25 Gly Arg Thr Met Asp Ser Ser Ala Glu Ser Lys Asp Arg Arg Glu Ala Gly Glu Asp Asn Glu Gly Phe Leu Ser Lys Leu Lys Lys Ile Phe Thr 25

Ser

<210> 26 <211> 6 <212> PRT <213> Homo sapiens <400> 26 Lys Arg Ser Thr Gly Asn <210> 27 <211> 6 <212> PRT <213> Mus sp.

<400> 27 Lys Arg Ser Thr Gly Asn 5